3. A method as recited in claim 1 wherein said promoter is a cauliflower mosaic virus promoter.

4. A method as recited in claim 1 wherein said promoter is an inducible promoter.

5. A method as recited in claim 1 wherein said promoter is a tissue specific promoter.--

In the specification:

In the specification following the title but before the first line of text, insert the following:

Cross-Reference to Related Applications

This is a divisional of Serial No. 09/638,715 , currently pending.

On page 3, lines 5-7, please amend the specification as follows:

--Accordingly, the object of this invention is to provide substrate-specific *DFR*s which have [altered] point mutations at residue number 134 of SEQ ID NO: 2 when the amino acids [sequences at the substrate specificity determining region] are aligned with the ClustalW program.--

On page 7, lines 7, please amend the specification as follows:

--using cDNA sequences of *Petunia* (SEQ ID NO: 35) and *Gerbera* (SEQ ID NO: 1).--

On page 13 after "Chimeric gene construction" please replace the present paragraph with the following paragraph:

--Highly conserved regions of the *DFR* gene were identified by a multiple sequence alignment of a number of DFRs. The 5' region (*Gerbera DFR* portion) of each chimeric gene was synthesized from the *Gerbera DFR* cDNA clone using a primer containing the codon for the starting methionine of the *Gerbera DFR* gene (SEQ ID NO. 5): [(]5'-GGC GAA AAT GGA AGA GGA TTC TCC-3'[)] and a primer containing a conserved region of the *Gerbera DFR* gene (Chimera 1; SEQ ID NO: 6: 5'-AGC AGA TGA AGT GAA CAC TAG TTT CTT CAC-3'; Chimera 2; SEQ ID NO: 7: 5'-GGC TTT CTC TGC